

A Comparative Study on Invasion, Survival, Modulation of Oxidative Burst, and Nitric Oxide Responses of Macrophages (HD11), and Systemic Infection in Chickens by Prevalent Poultry *Salmonella* Serovars

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Abstract

Poultry is a major reservoir for foodborne *Salmonella* serovars. *Salmonella* Typhimurium, *Salmonella* Enteritidis, *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg are the most prevalent serovars in U.S. poultry. Information concerning the interactions between different *Salmonella* species and host cells in poultry is lacking. In the present study, the above mentioned *Salmonella* serovars were examined for invasion, intracellular survival, and their ability to modulate oxidative burst and nitric oxide (NO) responses in chicken macrophage HD11 cells. All *Salmonella* serovars demonstrated similar capacity to invade HD11 cells. At 24 h post-infection, a 36–43% reduction of intracellular bacteria, in log₁₀(CFU), was observed for *Salmonella* Typhimurium, *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg, whereas a significantly lower reduction (16%) was observed for *Salmonella* Enteritidis, indicating its higher resistance to the killing by HD11 cells. Production of NO was completely diminished in HD11 cells infected with *Salmonella* Typhimurium and *Salmonella* Enteritidis, but remained intact when infected with *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg. Phorbol myristate acetate-stimulated oxidative burst in HD11 cells was greatly impaired after infection by each of the five serovars. When newly hatched chickens were challenged orally, a high rate (86–98%) of systemic infection (*Salmonella* positive in liver/spleen) was observed in birds challenged with *Salmonella* Typhimurium, *Salmonella* Enteritidis, *Salmonella* Heidelberg, and *Salmonella* Kentucky, while only 14% of the birds were *Salmonella* Senftenberg positive. However, there was no direct correlation between systemic infection and *in vitro* differential intracellular survival and modulation of NO response among the tested serovars.

Introduction

SALMONELLA ARE ONE OF THE LEADING causes of foodborne illness worldwide (Scallan *et al.*, 2011). In chickens, infections with host specific serovar *Salmonella* Gallinarum and *Salmonella* Pullorum cause septicemia fowl typhoid and pullorum disease, respectively (Barrow and Freitas Neto, 2011), whereas infections with non-host-specific serovars such as *Salmonella* Typhimurium, *Salmonella* Enteritidis, and *Salmonella* Heidelberg generally display no clinical symptoms. However, these non-host-specific poultry serovars account for the majority of clinical isolates in human salmonellosis. Although *Salmonella* Kentucky and *Salmonella* Senftenberg are not commonly associated with human salmonellosis, they, together with *Salmonella* Typhimurium, *Salmonella* Enteritidis,

and *Salmonella* Heidelberg, are the most common serovars isolated from U.S. poultry (CDC, 2008; FDA, 2010). *Salmonella* Senftenberg, a serovar that is more resistant to the environmental stresses, is frequently isolated from hatching houses and raw feed materials, and is adapted to colonize and persist in poultry houses (Liu *et al.*, 1969; Bailey *et al.*, 2001; Pedersena *et al.*, 2008). In the last decade, significant progress has been made in the knowledge of *Salmonella* invasion and pathogenesis in mammalian hosts, most of which are derived from studies based on the murine model of *Salmonella* Typhimurium infection (Haraga *et al.*, 2008; Malik-Kale *et al.*, 2011). Colonization by *Salmonella* in poultry has been extensively studied and well documented (Foley *et al.*, 2011); however, most are epidemiological investigations focused on prevalence. Information regarding *Salmonella* invasion and colonization mechanisms and

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interactions with host cells in chickens is limited and poorly defined (Boyd *et al.*, 2007; Lillehoj *et al.*, 2007; Chappell *et al.*, 2009; Wisner *et al.*, 2010, 2011).

Chicken macrophages play a critical role in the defense against microbial infection, in which they detect, phagocytize, and produce microbicidal substances, including reactive radical oxygen species (ROS), nitric oxide (NO), lysozyme, and proteolytic enzymes, to kill the infectious agents (Okamura *et al.*, 2005; Withanage *et al.*, 2005; Babu *et al.*, 2006). Robust macrophage functionality is associated with increased resistance to systemic spread (Wigley *et al.*, 2006) and intestinal colonization (Sun *et al.*, 2008) by *Salmonella*. However, the role of macrophages in controlling *Salmonella* infection and the interaction between *Salmonella* and macrophages in chickens are much less studied and remain mostly unclear.

In the present study, cell invasion, intracellular survival, and modulation of antimicrobial activity (NO and oxidative burst response) in chicken macrophage HD11 cells of the above mentioned five serovars were examined. Additionally, systemic infection by these five serovars in newly hatched chickens was also investigated.

Materials and Methods

Bacteria

Primary poultry isolates *Salmonella* Typhimurium and *Salmonella* Enteritidis were obtained from the National Veterinary Services Laboratory (Ames, IA) and were resistant to novobiocin–nalidixic acid (Kogut *et al.*, 1995). *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg used in this study were field isolates from broilers and were susceptible to novobiocin–nalidixic acid. All strains were susceptible to gentamicin. *Salmonella* from stocks were cultured overnight at 41°C in a Tryptic Soy Broth (TSB; Becton, Dickinson, and Company, Franklin Lakes, NJ) and the overnight cultures were transferred to a fresh TSB and cultured for 4 h to reach an exponential growth phase, and the bacteria were collected, washed, and resuspended in phosphate-buffered saline (PBS) at a final concentration of $\sim 2 \times 10^9$ colony-forming unit (CFU)/mL. Heat-killed *Salmonella* (HKS) were prepared by incubating the bacterial suspension at 75°C water bath for 15 min.

Chickens

Chickens (Hy-Line W36) were obtained from Hy-Line International (Bryan, TX) on the day-of-hatch. Birds were placed in floor pens with pine shavings in a controlled environment (biosafety level 2) and provided *ad libitum* access to water and a balanced unmedicated corn-soybean based diet with nutrient rations meeting or exceeding the recommendations of the National Research Council (NRC, 1994). The experiments comply with the Animal Care and Use Experimental Animal Protocol (Southern Plains Agricultural Research Center, Agriculture Research Service, U.S. Department of Agriculture).

HD11 Cells

The MC29 virus–transformed chicken macrophage cell line HD11 (Beug *et al.*, 1979) were maintained in complete Dulbecco's Modified Eagles Medium (DMEM; Invitrogen, Grand Island, NY) containing 10% chicken serum, antibiotics

(100 U penicillin/mL and 100 μ g streptomycin/mL), and 1.5 mM L-glutamine (Sigma, St. Louis, MO) at 39°C, 5% CO₂, and 95% humidity. Aliquots of cell suspension (2×10^6 cells/mL) were seeded into each well at 100 μ L/well for 96-well optical bottom black plate (Nalge Nunc International, Rochester, NY) and 500 μ L/well for 24-well plate (Becton Dickinson Biosciences, San Jose, CA) and allowed to grow to about 85% confluence (~ 36 h) before being used for assays. The 96-well plates were used for oxidative burst assay, and the 24-well plates were used for the cell invasion and the NO production assays.

Cell invasion and intracellular viability assay

Prior to infection, the culture medium was removed and cells were washed once and replaced with 200 μ L of plain DMEM (without chicken serum and other additives). Aliquots of 50 μ L of *Salmonella* suspensions ($\sim 2 \times 10^9$ CFU/mL) were added to each well with four replicate wells for each serovar and incubated for 1 h at 39°C in a 5% CO₂ humidified incubator. At 1 h post-infection (hpi), the infection medium was removed and the cells were washed once and treated with 100 μ g/mL of gentamicin sulfate in complete DMEM for 1 h to kill extracellular bacteria. After gentamicin treatment, infected cells were washed twice with PBS, lysed for 10 min in 300 μ L of 1% Triton X-100 (in PBS). After lysis, 700 μ L of PBS was added to each well and mixed thoroughly. Serial 1:10 dilutions were spread onto Difco's xylose-lysine tergitol 4 (XLT4) agar (Becton, Dickinson, and Company) plates and incubated at 41°C for 24 h. Colonies were counted to determine the CFU of intracellular bacteria at 2 hpi. Similarly, intracellular viable bacterial CFU was determined at 24 hpi after an additional culture of the infected cells for 22 h in the medium containing 20 μ g/mL of gentamicin sulfate.

NO production assay

Nitrite, a stable metabolite of NO, produced by activated macrophages was measured by the Greiss assay (Green *et al.*, 1982). HD11 cells in 24-well plates were treated, in four replicates, with live (prepared in the same way as above) or HKS as described in the killing assay. After 24 h of incubation, nitrite concentrations in the culture media were determined as previously described (He *et al.*, 2009).

Phorbol myristate acetate (PMA)–stimulated oxidative burst

Oxidative burst of HD11 cells was measured as described (He *et al.*, 2005). To evaluate the effect of *Salmonella* infection on oxidative burst of HD11 cells, PMA (Sigma) stimulated oxidative burst was measured in both *Salmonella*-infected and HKS-treated HD11 cells. The cells were first treated in plain DMEM with 10 μ L of live or HKS suspensions ($\sim 2 \times 10^9$ CFU) in a final volume of 50 μ L/well at 39°C for 1 h. Following the treatment, the cells were washed and stimulated in plain DMEM with PMA (0.5 μ g/mL) in a final volume of 100 μ L/well containing 10 μ g/mL of 2',7'-dichlorfluorescein-diacetate (DCFH-DA; Sigma) and 100 μ g/mL of gentamicin sulfate for 1 h at 39°C in 5% CO₂ and 95% humidity. The relative fluorescent units (RFU) at the end of incubation were measured (485/530 nm) using Genios Plus Plate Reader (Tecan US Inc., Durham, NC).

In vivo organ invasion

Day-old chickens, 25 per group, were orally challenged with 0.5 mL of each different *Salmonella* serovar ($\sim 5 \times 10^8$ CFU/bird) and housed in separated rooms. At 4 days post-infection (dpi), chickens were euthanized with CO₂, and liver and spleen were aseptically removed from each chicken and cultured as a combined sample in tetrathionate broth overnight (18–24 h) at 41°C according to guidelines of the U.S. Department of Agriculture (USDA, 1989). After incubation, aliquots of 10 μ L of broth were streaked on XLT4 plates and incubated for 24 h at 41°C. Two independent experiments were conducted at different dates, and a total of 50 chickens were used for each treatment group.

Statistical analysis

At least three independent experiments for NO, oxidative burst, and invasion, and two separate experiments for organ invasion were conducted. Statistical difference was determined at the level of $p < 0.05$ by Student's *t*-test using SigmaStat software (Jandel Corp., Richmond, CA).

Results

Differential modulation of NO production in HD11 cells by *Salmonella* serovars

Salmonella infection-induced NO production in HD11 cells was serovar-dependent (Fig. 1A): *Salmonella* Typhimurium and *Salmonella* Enteritidis induced little or no output of NO, while *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg stimulated significant amounts of NO production. This inhibition of NO production in HD11 cells was observed only in treatments with live *Salmonella* Typhimurium and *Salmonella* Enteritidis; HD11 cells stimulated with HKS showed a strong NO production regardless of the *Salmonella* serovar (Fig. 1B).

Down-regulation of HD11 cell oxidative burst potential by intracellular *Salmonella*

To examine the effect of intracellular *Salmonella* on the oxidative burst capacity of HD11 cells, PMA was used to stimulate oxidative burst in HD11 cells infected with *Salmonella* or treated with HKS. After invasion, all *Salmonella* serovars tested significantly diminished the oxidative burst potential of HD11 cells and rendered the macrophages irresponsive to PMA stimulation (Fig. 2A). Metabolically inhibition by intracellular *Salmonella* was the most likely cause for the loss of oxidative response of *Salmonella*-infected cells to PMA stimulation, since HD11 cells pretreated with HKS in an identical manner displayed no inhibitory effect on oxidative burst response to a subsequent stimulation with PMA (Fig. 2B).

Cell invasion and intracellular survival in HD11 cells by *Salmonella*

All *Salmonella* serovars demonstrated similar capacity ($p \geq 0.05$) to invade HD11 cells as demonstrated by the CFU at 2 hpi, even though *Salmonella* Typhimurium and *Salmonella* Enteritidis invasion was numerically greater than *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg (Table 1). The viability of *Salmonella* Enteritidis at 24 hpi, however, was significantly higher ($p < 0.05$) than that of

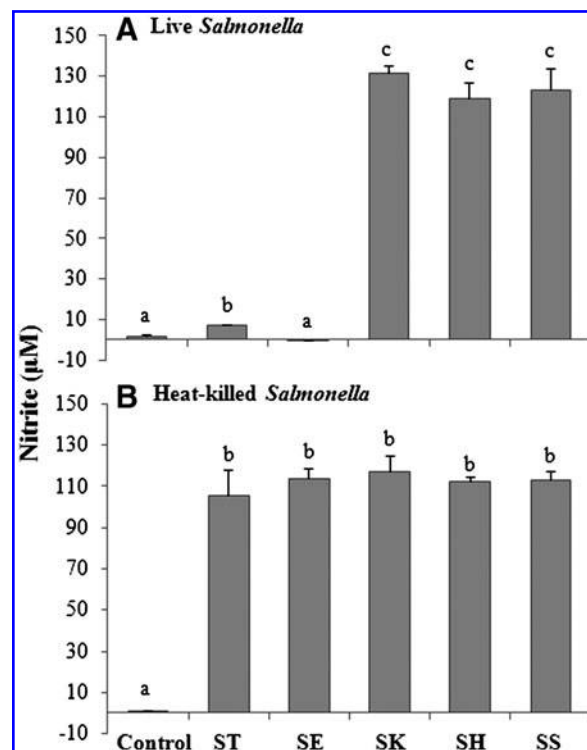


FIG. 1. Effect of *Salmonella* infection on nitric oxide (NO) production in HD11 cells. HD11 cells were infected with *Salmonella* for 1 h in 24-well plates at 39°C in a 5% CO₂ humidified incubator. At 1 h post-infection (hpi), extracellular *Salmonella* were killed by incubation with media containing 100 μ g/mL of gentamicin sulfate for 1 h; the cells were washed and then cultured for an additional 22 h in a medium containing 20 μ g/mL of gentamicin sulfate; and nitrite contents in cell culture media were determined. Treatment with heat-killed *Salmonella* (HKS) was performed identically as with live *Salmonella*. (A) HD11 infected with live *Salmonella*. (B) HD11 treated with HKS. ST, *Salmonella* Typhimurium; SE, *Salmonella* Enteritidis; SK, *Salmonella* Kentucky; SH, *Salmonella* Heidelberg; SS, *Salmonella* Senftenberg. Different letters indicate that the difference between these groups is statistically significant ($p < 0.05$).

Salmonella Typhimurium, *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg, indicating that *Salmonella* Enteritidis was the most resistant to intracellular killing by HD11 cells among the five serovars.

Systemic invasion by *Salmonella* in neonatal chickens

Systemic infection in young chickens by these five *Salmonella* serovars was investigated by examining the presence of *Salmonella* in the liver/spleen of challenged birds at 4 dpi (Table 2). A high percentage (86–98%) of positive birds was observed in groups infected with *Salmonella* Typhimurium, *Salmonella* Enteritidis, *Salmonella* Heidelberg, and *Salmonella* Kentucky, whereas only 14% of the birds were positive in the *Salmonella* Senftenberg challenged group. The unchallenged control birds were all *Salmonella* negative.

Discussion

Survival inside the macrophage is essential for *Salmonella* virulence and systemic infection (Fields *et al.*, 1986; Schwan

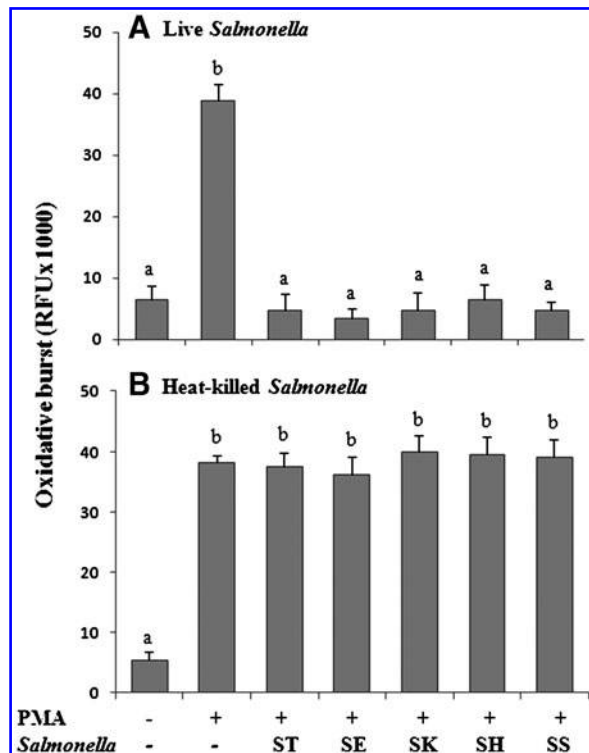


FIG. 2. Effect of *Salmonella* infection on Phorbol myristate acetate (PMA)-stimulated oxidative burst in HD11 cells. HD11 cells were infected with *Salmonella* or treated with heat-killed *Salmonella* (HKS) for 1 h in 96-well optical bottom black plates at 39°C in a 5% CO₂ humidified incubator. At 1 h post-infection (hpi), the cells were washed and stimulated with PMA (0.5 µg/mL) in a final volume of 100 µL/well containing 10 µg/mL of DCFH-DA and 100 µg/mL of gentamicin sulfate for 1 h. The relative fluorescent units (RFU) at the end of incubation were measured (485/530 nm). (A) HD11 infected with live *Salmonella*. (B) HD11 treated with HKS. ST, *Salmonella* Typhimurium; SE, *Salmonella* Enteritidis; SK, *Salmonella* Kentucky; SH, *Salmonella* Heidelberg; SS, *Salmonella* Senftenberg. Different letters indicate that the difference between the groups is statistically significant ($p < 0.05$).

TABLE 1. INTRACELLULAR SURVIVAL OF *SALMONELLA* IN HD11 CELLS^a

Serovar	CFU (2 hpi)	CFU (24 hpi)	Reduction (%)
<i>Salmonella</i> Typhimurium	6.52 ± 0.47	3.63 ^b ± 0.40	43.29 ^b ± 3.79
<i>Salmonella</i> Enteritidis	6.63 ± 0.44	5.60 ^c ± 0.25	16.40 ^c ± 6.94
<i>Salmonella</i> Kentucky	5.85 ± 0.51	3.74 ^b ± 0.78	35.98 ^b ± 13.32
<i>Salmonella</i> Heidelberg	5.93 ± 0.43	3.66 ^b ± 0.46	37.92 ^b ± 4.68
<i>Salmonella</i> Senftenberg	5.95 ± 0.35	3.79 ^b ± 0.61	35.70 ^b ± 8.00

^aHD11 cells were infected with *Salmonella* at 2×10^9 CFU/mL for 1 h at 39°C in a 5% CO₂ humidified incubator and intracellular viable *Salmonella* (CFU in log₁₀ scale) at 2 and 24 h post-infection (hpi) were counted. Data are mean ± standard deviations of CFU/well. Reduction (%) = [(CFU (2 hpi) - CFU (24 hpi)) / CFU (2 hpi)] / 100. Different letters indicate that the differences between these groups are statistically significant ($p < 0.05$).

TABLE 2. ORGAN INVASION BY *SALMONELLA* IN NEWLY HATCHED CHICKENS^a

Serovar	Trial 1	Trial 2	Mean ± SD
<i>Salmonella</i> Typhimurium	19/25	24/25	86 ± 14
<i>Salmonella</i> Enteritidis	25/25	23/25	96 ± 6
<i>Salmonella</i> Kentucky	24/25	25/25	98 ± 3
<i>Salmonella</i> Heidelberg	24/25	24/25	96 ± 0
<i>Salmonella</i> Senftenberg	3/25	4/25	14 ± 3

^aDay-old chickens, 25 per group, were orally challenged with different *Salmonella* serovar ($\sim 5 \times 10^8$ CFU/bird). At 4 days post-infection (dpi), chickens were euthanized, and liver and spleen were removed to test for organ invasion by *Salmonella*. Data in columns for Trials 1 and 2 are *Salmonella*-positive birds in each group at 4 dpi. Data in column for mean ± SD are means and standard deviations of the *Salmonella*-positive birds (%) in each group of the two trials.

et al., 2000; Guiney, 2005). *Salmonella* virulence depends at least partially on the type III secretion system (T3SS), which secretes and delivers nearly 40 different virulence effectors into host cells, to facilitate invading, surviving, and replicating within host cells (Haraga *et al.*, 2008; Ibarra *et al.*, 2009; Malik-Kale *et al.*, 2011). To defend against the host cell antimicrobial defense mechanisms, *Salmonella* produce effector proteins which manipulate host cells to delay the phagolysosomal maturation and hence avoid exposure to lysosomal contents (Haraga *et al.*, 2008) and secrete various metabolic enzymes which neutralize the antimicrobial effect of free radicals oxygen and nitrogen species (ROS and RNS) (Aussel *et al.*, 2011; Henard and Vázquez-Torres, 2011; Schlauch, 2011).

Professional phagocytes generate ROS in the process of an oxidative burst during phagocytosis of microbes or in response to stimulation by microbial components (Fang, 2011). ROS production in response to microbe and microbial component stimulation play a critical role in controlling microbial infection (Ogier-Denis *et al.*, 2008; Lam *et al.*, 2010). The exact role of ROS in controlling intracellular *Salmonella* in macrophages is debatable (Fang, 2011), since *Salmonella* carry abundant enzymes (catalases, peroxiredoxins, superoxide dismutases) to neutralize the effect of ROS (Aussel *et al.*, 2011; Fang, 2011). PMA is a protein kinase C activator and stimulates a strong oxidative burst in chicken phagocytes (He *et al.*, 2005). The effect of intracellular *Salmonella* on the oxidative burst response of HD11 cells to PMA stimulation has not been reported and therefore was examined in the present study. Macrophage HD11 cells infected with the five *Salmonella* serovars showed severely impaired ROS response to PMA stimulation as compared to the non-infected cells, while the cells treated identically with dead *Salmonella* (HKS) demonstrated a normal ROS response to PMA. The lack of ROS response to PMA stimulation in *Salmonella*-infected HD11 cells is likely caused by intracellular *Salmonella* which may inhibit phagocyte NADPH oxidase activity and metabolically neutralize the ROS products. Our results clearly indicate that *Salmonella* serovars are well adapted to evade the ROS-mediated killing in macrophage. Although the ROS-mediated direct killing of intracellular *Salmonella* might be limited in chicken macrophages as our results suggest, accumulated evidence suggests that ROS can act as signaling molecules to indirectly assert an antimicrobial role. For example, ROS have been reported to activate MAP kinase and transcription factors NF-κB and AP-1, up-regulate inflammatory cytokine and

chemokine expression, and induce the formation of autophagy (Torres and Forman, 2003; Closa and Folch-Puy, 2004; Huang *et al.*, 2011). Activation of these cellular functions plays a critical role in controlling intracellular *Salmonella* (Rosenberger and Finlay, 2002; Sahlberg *et al.*, 2007; Jones *et al.*, 2008; Deretic, 2011).

NO response to microbial stimulation is an important innate immune function of macrophages and plays a critical role in controlling the proliferation of intracellular bacterial pathogens such as *Salmonella* Typhimurium (Mastroeni *et al.*, 2000; Alam *et al.*, 2002; 2008). However, virulent factors secreted via *Salmonella* T3SS can suppress iNOS activity (Das *et al.*, 2009) and prevent iNOS-containing vesicle trafficking to phagosomes, hence limiting exposure of *Salmonella* to RNS (Chakravorty *et al.*, 2002). Additionally, *Salmonella* possess three major enzymes (flavo-hemoglobin Hmp, flavorubredoxin NorV, and cytochrome *c* nitrite reductase NrfA) that can detoxify NO under different environmental conditions (Bang *et al.*, 2006; Mills *et al.*, 2008). Previously, infection with *Salmonella* Typhimurium and *Salmonella* Enteritidis has been shown to induce NO production in chicken macrophages (Okamura *et al.*, 2005; Withanage *et al.*, 2005; Babu *et al.*, 2006). However, our results show that NO production in HD11 cells was completely inhibited by infection with *Salmonella* Enteritidis and only minor amounts of NO was produced in cells infected with *Salmonella* Typhimurium. Infection of HD11 cells with *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg induced large amounts of NO. The results demonstrate a dramatic difference in their ability to modulate host cell NO response among these wild-type *Salmonella* serovars. The lack of or diminished NO response in HD11 cells to *Salmonella* Typhimurium and *Salmonella* Enteritidis infection is probably due to the inhibition on iNOS or the metabolic detoxification of NO, since heat-killed *Salmonella* Typhimurium and *Salmonella* Enteritidis were able to induce large quantities of NO comparable to the levels induced by the other three serovars. The exact mechanism that enables *Salmonella* Typhimurium and *Salmonella* Enteritidis to prevent NO production in HD11 cells is not clear and needs to be further investigated. The discrepancy in NO induction by *Salmonella* Typhimurium and *Salmonella* Enteritidis infection between this study and previous studies (Okamura *et al.*, 2005; Withanage *et al.*, 2005; Babu *et al.*, 2006) cannot be readily explained and may be due to different conditions under which these experiments were conducted.

There was no difference in the rate of internalization of *Salmonella* among the serovars used in this study. Within 24 hpi, HD11 cells were able to limit intracellular *Salmonella* growth and achieved reduction of viable intracellular *Salmonella* of all serovars tested. However, *Salmonella* Enteritidis had the highest viability at 24 hpi among the tested serovars, indicating that *Salmonella* Enteritidis was the most resistant against macrophage-mediated bacterial killing. These results provide supporting evidence to an epidemiological observation that *Salmonella* Enteritidis is more adapted to cause systemic infections in chickens (Foley *et al.*, 2011). There was no apparent indication that the ability of *Salmonella* to down-regulate ROS and RNS responses provided an advantage in survival within chicken macrophages, since *Salmonella* Typhimurium had a similar, if not the lowest, viability among the five tested serovars despite the fact that it almost completely abrogated NO production in HD11 cells. Therefore,

our results indicate that direct killing by ROS or RNS may not play a determinant role in intracellular survival of *Salmonella* in chicken macrophages.

The ability of *Salmonella* to invade *via* the intestine and spread systemically plays an important role in *Salmonella* colonization of reproductive organs, a main cause for internal contamination of eggs (Gast *et al.*, 2004). Systemic infection and reproductive organ colonization by serovars *Salmonella* Enteritidis, *Salmonella* Typhimurium, and *Salmonella* Heidelberg has been previously reported (Gantois *et al.*, 2008). However, the present study was the first to report that *Salmonella* Kentucky is capable of causing systemic infection in liver and spleen through intestinal invasion in young chickens. It is interesting to note that chickens challenged with *Salmonella* Senftenberg had an exceptionally low rate (14%) of systemic infection as compared to chickens challenged with other serovars (86–98%). Although *Salmonella* Senftenberg has been reported to persist in and frequently isolated from poultry hatching houses, farm houses, and raw feed materials (Liu *et al.*, 1969; Bailey *et al.*, 2001; Kim *et al.*, 2007; Pedersen *et al.*, 2008), it remains a less prevalent strain in chicks, hens, and poultry products. Our results demonstrated that *Salmonella* Senftenberg lacks the ability to attain systemic infection, suggesting this strain is deficient in its ability to invade.

In summary, we have examined the five most prevalent *Salmonella* serovars in U.S. poultry for intracellular survival and their ability to modulate antimicrobial activity in chicken macrophage HD11 cells. Intracellular *Salmonella* impaired the oxidative burst response of HD11 to PMA stimulation. Infection with *Salmonella* Typhimurium and *Salmonella* Enteritidis, but not *Salmonella* Heidelberg, *Salmonella* Kentucky, and *Salmonella* Senftenberg, abolished NO production in HD11 cells. Serovar *Salmonella* Enteritidis was best adapted to survive inside HD11 cells among the tested serovars. Newly hatched chickens were vulnerable to systemic infection by *Salmonella* Typhimurium, *Salmonella* Enteritidis, *Salmonella* Heidelberg, and *Salmonella* Kentucky, but not *Salmonella* Senftenberg. These results demonstrate the different capabilities to modulate the immune response in chicken macrophages among *Salmonella* serovars; however, further investigations are needed to identify factors that control intracellular survival and systemic infection.

Disclosure Statement

No competing financial interests exist.

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